	(X1) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
CCC	CGATCGGC CGCATGC	1
(2)	INFORMATION FOR SEQ ID NO:16:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 51 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
GCA	TTTAAAG AATGGGAAGG GATCCTAGGA ATCCAGCAAC CAGGACCAGA G	51
(2)	INFORMATION FOR SEQ ID NO:17:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
GAG	CTCTTGT TAAAAAAGGT GTTCCAGATC	30
(2)	INFORMATION FOR SEQ ID NO:18:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 62 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ix) FEATURE: (A) NAME/KEY: modified_base (B) LOCATION: 1939 (D) OTHER INFORMATION: /note= "N = G, A, T or C"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
	TOTAL AND COMMAND AND AND AND AND AND AND AND AND AND	60
GCA	TTTAAAG AATGGGAANN NNNNNNNNN NNNNNNNNA CCAGGACCAG AGTAATTGAT	•

(2) INFORMATION FOR SEQ ID NO:19:

5	(1) SEQUENCE CHARACTERISTICS: (A) LENGTH: 55 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
10	GGGCTACTTG AAAGGGACAT TCCTTCGTTT GCAATTTCTG GATTTGAAGT ACCCC	55
	(2) INFORMATION FOR SEQ ID NO:20:	
15	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
20 25 25	CCAAGAAAT ACTAGAGCTC TTGTTAAAAA AGGTGTTCC	39
25	(2) INFORMATION FOR SEQ ID NO:21:	
30	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
35	GAGATTCTGT AATTTTTGGA GAAGCATGGG GGTTGACAAC GATAAATGTC	50
	(2) INFORMATION FOR SEQ ID NO:22:	
40 45	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 63 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
73	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
	GCATTTAAAG AATGGGAAGA AGATCCTAAT AATCCAGCAA CCAGGACCAG AGTAATTGAT	60
50	CGC	63
	(2) INFORMATION FOR SEQ ID NO:23:	

	(I) SEQUENCE CHARACTERISITES:	
	(A) LENGTH: 7 amino acids	
	(B) TYPE: amino acid	
	(C) STRANDEDNESS: single	
5	(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
	Glu Asp Pro Asn Asn Pro Ala	
10	1 5	
	(2) INFORMATION FOR SEQ ID NO:24:	
15	(i) SECTIONS CHARACTERISTICS.	
13	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 51 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
20 25		
T	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
	GCATTTAAAG AATGGGAAGG GATCCTAGGA ATCCAGCAAC CAGGACCAGA G	51
25		
nete:	(2) INFORMATION FOR SEQ ID NO:25:	
11	(i) SEQUENCE CHARACTERISTICS:	
:	(A) LENGTH: 63 base pairs	
3 0	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
ine in	(b) Torobodi. Timeat	
of it Design	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
3 5	(XI) DEGULACE DESCRIPTION. SEQ ID NO.25.	
	GCATTTAAAG AATGGGAAGA TGATCCTCAT AATCCCACAA CCAGGACCAG AGTAATTGAT	60
		0.0
	CGC	63
40		
	(2) INFORMATION FOR SEQ ID NO:26:	
	· · · · · · · · · · · · · · · · · · ·	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 7 amino acids	
45	(B) TYPE: amino acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(b) TOPOLOGI. TIMEAT	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
50	(AI) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
20	Agn Agn Dro Hig Agn Dro Thr	
	Asp Asp Pro His Asn Pro Thr	
	i	